

cell proliferation⁺--;

line 8, please delete "proliferation⁺";

line 10, please amend the second to last column to read --25-- and the last column to read --102--.

IN THE CLAIMS:

Please cancel claims 14-16 and 18 from the application, without prejudice. Please amend claims 1, 3, 7, 17, 19, 20 and 21 as follows:

- 1: (Twice amended) A method for making [at least a portion of] a humanized antibody variable domain comprising amino acid sequences of an import antibody comprising a non-human antibody which is desired to be humanized [(import antibody)] and a human antibody, comprising the steps of:
- a. obtaining the amino acid sequences of an import variable domain and of a consensus human variable domain of a human immunoglobulin subgroup *New*
 - b. identifying Complementarity Determining Region (CDR) amino acid sequences in the import and the human amino variable domain sequences;
 - c. substituting an import CDR amino acid sequence for the corresponding human CDR amino acid sequence;
 - d. aligning the amino acid sequences of a Framework Region (FR) of the import antibody and the corresponding FR of the consensus antibody;
 - e. identifying import antibody FR residues in the aligned FR sequences that are non-homologous to the corresponding consensus antibody residues;
 - f. determining if the non-homologous import amino acid residue is [reasonably] expected to have at least one of the following effects:
 1. non-covalently binds antigen directly,
 2. interacts with a CDR; or
 3. participates in the V_L - V_H interface by affecting the proximity or orientation of the V_L and V_H regions with respect to one another;